

Sequence listings

SEQ ID NO: 1

Homo sapiens transforming growth factor, beta receptor II (70/80kDa),
complete antisense sequence

Derived from AY675319

1 CAGCCACACT GTCTTTAAGT CTCAGCCCAC CCACACTGAG GAGGGTGCCT AGAGGTTCTA
61 TTTCCAAACC TTTGCATGTA TCTTAAAAAT CTCAATAAAA TGAGACCTTC CACCATCCAA
121 ACAGAGCTGA TATTCTCACT ACCAGTCCCT CTCTAATATT CCTATTTGGC TGAAAATAAG
181 TAGCTTCAAA AAGTTTAAA AAAGAGATTA CTTGCAGCAT TAACACTTCT TTGTTGATTA
241 ACAAGTTCC TATGGAGTT TAAAGCTCAT ACTTTGTTCT TGTCCCTGTG GACACAAATT
301 TTCTAACTGC AAATGGGACC TTTGTGTCCC ACATTCAAAT CCTCTCTAGT AATTTCTGCA
361 AAGGTTGAGA AGGCTGGCAT GATGGAGAGA ACGGTAACCA TGAGGAAAGC TTCTTGGAGT
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481 GGGCTAACTG AGACCCTTAA AGGAGTTCCC CTTAGTCCA ATAAAAGGCC AACCTCAAAT
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601 GGACATTCT TACAAGGGAC CTTGGTTAGG TGCAGATTAA ATTCCCTAGAC TGGGGTCCAG
661 GTAGGCAGTG GAAAGAGCTA ATGTTACAG TGAGAAGTGA GGCAGCTTG TAAGTGTCTC
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Homo sapiens transforming growth factor, beta receptor II (70/80kDa), antisense sequence

Derived from BC040499

1 tttttttttt tttttctag gaatggaaac aggaggcagg atgctcacct gagtatttg
61 ctttattcaa tctaataaaac attttattta tgtaaaagac aaacaatgca tagaataaaa
121 ataagtgctt gagactttg atataaaaag agtataatgc attcacattc ctatTTaat
181 acatgagtagc agctgaagtg ttccataaaaa gaataaaaact ttccctttat gtatagtagt
241 gaaaaaaagtc agtattttt ggaactacag aatgttattc cttggttttt tttcttgaat
301 aagaaaaaaa aacataaaaca aaacaagcca cagtatcctc tgacactaca ttccagttt
361 tgctgataac ccagaagtga gaatactctt gaatcttcaa tatctcatga atggaccagt
421 attctagaaa ctcaccacta gaggtcaatg ggcaacagct attggatgg tatcagcatg
481 ccctacggtg caagtggaaat ttctaggcgc ctctatgcta ctgcagccac actgtcttt
541 actctcagcc cacccacact gaggagggtg cctagaggtt ctatTTcaa acctttgcatt
601 gtatcttaaa aatctcaata aatgagacc ttccaccatc caaacagagc tgatattctc
661 actaccagtc cctctctaattt attccttattt ggctgaaaat aagtagcttc aaaaagttt
721 aaaaaagaga ttacttgcag cattaacact tctttgttga ttaacaagtt tcctatggag
781 ttttaaagct catactttgt tcttgcctt gtggacacaa atttcttaac tgcaaatggg
841 acctttgtgt cccacattca aatcctctct agtaatttct gcaaagggtg agaaggctgg
901 catgatggag agaacggtaa ccatgaggaa agcttcttgg agtaaagcac tcctctctcc
961 aatgcagagg gtaaaaactat taacatataa gcaaaagaaa cttgggctaa ctgagaccct
1021 taaaggagtt ccccttagt ccaataaaag gccaacttca aatcttaaca ccagataagg
1081 tagtcaaaat catattatat acccagagaa tgactgcttgc aatggacatt tcttacaagg
1141 gaccttggtt aggtgcagat ttaattccta gactgggtcc aggtaggcag tggaaagagc
1201 taatgttac agtgagaagt gaggcagtt tgtaagtgtc tccacacctt cacattttgt
1261 gaacgtggac tggagataac tgaaaaccat ctgctatcct tacctgggaa tccagatttt
1321 cctgcaaaat ctccaaatat ttataaagtg gcttcacttt ttgaaacgct gtgctgacca
1381 aacaaaacat atgttagag tgcctgaggt catagtcctg acaatgatag tattgtgttag
1441 ttgaaatcct cttcatcagg ccaaactgtg cttgagcaat caggagccca gaaagatgg
1501 acccatttgtt gttgtatag aaaactagaa aatcaagtca agttaatga aaaaagtaaac
1561 acgataaaagc ctagagttag aatttgcctt ttttttagaaa agatgaagg ctgggagcag
1621 agaatagtaa cataagtgcata gggaaagat gaaaaaaaaga acaatTTTC attagtagat
1681 ggtggggcaa tcgcattggat gggacatct gttctgattt ttctgcaacc catgaaggta
1741 aaaaagtgggg ttcaaaaacat tcaaggattt aaagatgggg tagatTTCT aaacttaggtt
1801 gagggagagt ttctaaacta gccccccaga tttggggctt ggagctaaa tgaaaagtcc
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1921 ctttagttcca tggccagaag agaagtgcata ggcaggaaat gatttttg caaaagcaag
1981 tgcaatgtgg tcatacgctgg ctgtgagaca tggagcctt ttcctcatgc aaagttcact
2041 gttttacagt cagagaacca ctgcattgtgt gattgtcaaa tgctaatgct gtcattggc
2101 ctttccttct ctgcttggtt ctggagttct ccaataaaac caatttcctg ggaatatttg
2161 atgttttcc ttgtctttt tcaaggatg gctatataa tagagctata gacatataa
2221 gatatatata tatatatata aaacatagct attcatattt atatacaggc attaataaaag
2281 tgcaaatgtt attggctatt gtaaaaatca atctcatttc ctgaggaagt gctaacacag

2341 cttatcctat gacaatgtca aaggcataga atgctctatg tcacccactc cctgctgctg
2401 ttgtttctgc ttatccccac agcttacagg gaggggagtg accccccttgg tttccagga
2461 agcatcagtt caggggcagc ttccctgctgc ctctgttctt tggtgagagg gcagcctctt
2521 tggacatggc ccagcctgcc ccagaagagc tatttggtag tgtttaggga gccgtcttca
2581 ggaatcttct cctccgagca gctccccc gagagcctgt ccagatgctc cagctcactg
2641 aagcgttctg ccacacactg ggctgtgaga cgggcctctg ggtcgtggc ccagcactca
2701 gtcaacgtct cacacaccat ctggatgccc tgggtggtaa gccagaagct gggaaattct
2761 ggtcgccctc gatctctcaa cacgttgc ttcatgctt cgacacaggg gtgctccgc
2821 accttggAAC caaatggagg ctcataatct tttacttctc ccactgcatt acagcgagat
2881 gtcattttccc agagcaccag agccatggag tagacatcg tctgcttggaa ggactcaaca
2941 ttctccaaat tcatcctgga ttcttaggact tctggagcca tgtatcttgc agttcccacc
3001 tgcccactgt tagccaggc atccacagac agagtagggg ccagacgcag gaaaagccca
3061 aagtcacaca ggcagcagg taggtcggtc ttcacgagga tattggagct cttgagggtcc
3121 ctgtgcacga tggcatctt gggctccca catggagtgt gatcactgtg gaggtgagca
3181 atccccccggg cgagggagct gcccagcttgc cgccaggtcct cccagctgat gacatgccgc
3241 gtcaggtact actgttagggtt gcccctggcg tggaaggcgg tgcgttgcgttgc gtattgttgc
3301 cccaaactccg tcttccgctc ctcagccgtc aggaactgga gtatgttctc atgcttcaga
3361 ttgatgtctg agaagatgtc cttctctgtc ttccaagagg catactcctc atagggaaag
3421 atcttgactg ccactgtctc aaactgctct gaagtgttct gcttcagctt ggccttata
3481 acctcagcaa agcgacctt ccccaccagg gtgtccagct caatggcag cagctctgt
3541 ttgtggtaa tttgtggc acacgtggag ctgtatgtc agcggtcatttcc ttccaggatg
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3661 ctcagcttct gtcgttgc aacgcggtag cagtagaaga tgcgtatgac agatatggca
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3781 ggattgtgg ttttatattt ttctgttgc atgatgttgc cattgcactc atcagagct
3841 caggaacaca tgaagaaagt ctcaccaggc tttttttt ctttcataat gcactttgg
3901 gaagcagcat cttccagaat aaagtcatgg taggggagct tgggttgc gcaactgtc
3961 tcttagtta tttctcggtc attcttctc catacagcca cacagacttc ctgtggcttc
4021 tcacagatgg aggtgtatgtc gcagttgtc atgcaggatt tctgggttgc acaggtggaa
4081 aatctcacat cacaatattt acacgttgc ggaaacttgc ctgcaccgtt gttgtcagt
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4201 gtccacagga cgatgtgcag cggccacagg cccctgagca gccccggacc catggcagac
4261 cccgctgtc gtcgttgc gaccccccgg cgcagcggac ggcgccttcc cggaccctg
4321 gctgcgcctc cgcgcgcgc cctctccggc ccccgccgg ggcggcagc gcagatgtgc
4381 gggccagatg tggcgccgc tcgttgcggc ggagggggcc tggaggccgg cgaggcgcgg
4441 ggaggcccccc ggcggccgg ggaagctgca caggagttcg gtcctgtcc cgagcgggt
4501 cacgcgcggg ggtgtcggtc ctccgtgcgc gcgagtgact cactcaactt caactcagcg
4561 ctgcggggaa aacagggaaac tcctcgccaa cagctggca ggacctctct ccgccccgaga
4621 gccttctccc tctccaa

SEQ ID NO: 3

CAGCCCCCGACCCATG

DNA

Artificial Sequence

SEQ ID NO: 4

GCTGATGCCCTGTCACTTGAA

DNA

Artificial Sequence

SEQ ID NO: 5

GCCATGGAGTAGACATCGGT

DNA

Artificial Sequence

SEQ ID NO: 6

GCAACAGCTATTGGGATGGT

DNA

Artificial Sequence

SEQ ID NO: 7

GTGCAGGGAAAGATGAAAA

DNA

Artificial Sequence

SEQ ID NO: 8

GTATCAGCATGCCCTACGGT

DNA

Artificial Sequence

SEQ ID NO: 9

GGATCCAGATTTCTGCCTAA

DNA

Artificial Sequence

SEQ ID NO: 10

GGAGAAGCAGCATCTTCCAG

DNA

Artificial Sequence

SEQ ID NO: 11
GAGCTCTTGAGGTCCCTGTG
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Artificial Sequence

SEQ ID NO: 12
GAGACCTTCCACCATCCAAA
DNA
Artificial Sequence

SEQ ID NO: 13
TAGCTGGCTGTGAGACATGG
DNA
Artificial Sequence

SEQ ID NO: 14
TTTGAAACGCTGTGCTGAC
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Artificial Sequence

SEQ ID NO: 15
TCAGCCAGTATTGTTCCCC
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Artificial Sequence

SEQ ID NO: 16
TCACACAGGCAGCAGGTTAG
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SEQ ID NO: 17
TCAGGAATCTTCTCCTCCGA
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Artificial Sequence

SEQ ID NO: 18

TGGTAGTGTAGGGAGCCG

DNA

Artificial Sequence

SEQ ID NO: 19

TATCCCCACAGCTTACAGGG

DNA

Artificial Sequence

SEQ ID NO: 20

AGCCTCTTCCTCATGCAA

DNA

Artificial Sequence

SEQ ID NO: 21

ATGTCATTTCCCAGAGCACC

DNA

Artificial Sequence

SEQ ID NO: 22

AGGAATCTTCTCCTCCGAGC

DNA

Artificial Sequence

SEQ ID NO: 23

AGCCATGGAGTAGACATCGG

DNA

Artificial Sequence

SEQ ID NO: 24

ATGCTACTGCAGCCACACTG

DNA

Artificial Sequence

SEQ ID NO: 25

CCTTCTCTGCTTGGTTCTGG

DNA

Artificial Sequence

SEQ ID NO: 26

CCAGGAGAAATAAGGGCACA
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Artificial Sequence

SEQ ID NO: 27
CAGCAGCTCTGTGTTGTGGT
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Artificial Sequence

SEQ ID NO: 28
CCCACTGTTAGCCAGGTCAT
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Artificial Sequence

SEQ ID NO: 29
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SEQ ID NO: 31
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Artificial Sequence

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Artificial Sequence

SEQ ID NO: 37

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Artificial Sequence

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Artificial Sequence

SEQ ID NO: 41

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Artificial Sequence

SEQ ID NO: 42

GCAGCCCCCGACCCATGGC

DNA

Artificial Sequence

SEQ ID NO: 43

GCAGCCCCCGACCCATGG

DNA

Artificial Sequence

SEQ ID NO: 44

GCAGCCCCCGACCCATG

DNA

Artificial Sequence

SEQ ID NO: 45

AGCAGCCCCCGACCCATGGCAGAC

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Artificial Sequence

SEQ ID NO: 46

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DNA

Artificial Sequence

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Artificial Sequence

SEQ ID NO: 48

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Artificial Sequence

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Artificial Sequence

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AGCAGCCCCGACCCATGG
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Artificial Sequence

SEQ ID NO: 51
AGCAGCCCCGACCCATG
DNA
Artificial Sequence

SEQ ID NO: 52
GAGCAGCCCCGACCCATGGCAGA
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SEQ ID NO: 53
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SEQ ID NO: 54
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SEQ ID NO: 55
GAGCAGCCCCGACCCATGGC
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Artificial Sequence

SEQ ID NO: 56

GAGCAGCCCCGACCCATGG

DNA

Artificial Sequence

SEQ ID NO: 57

GAGCAGCCCCGACCCATG

DNA

Artificial Sequence

SEQ ID NO: 58

TGAGCAGCCCCGACCCATGGCAG

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Artificial Sequence

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Artificial Sequence

SEQ ID NO: 60

TGAGCAGCCCCGACCCATGGC

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Artificial Sequence

SEQ ID NO: 61

TGAGCAGCCCCGACCCATGG

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Artificial Sequence

SEQ ID NO: 62

TGAGCAGCCCCGACCCATG

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Artificial Sequence

SEQ ID NO: 63

CTGAGCAGCCCCGACCCATGGCA

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Artificial Sequence

SEQ ID NO: 64

CTGAGCAGCCCCGACCCATGGC

DNA

Artificial Sequence

SEQ ID NO: 65

CTGAGCAGCCCCGACCCATGG

DNA

Artificial Sequence

SEQ ID NO: 66

CTGAGCAGCCCCGACCCATG

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Artificial Sequence

SEQ ID NO: 67

CCTGAGCAGCCCCGACCCATGG

DNA

Artificial Sequence

SEQ ID NO: 68

CCTGAGCAGCCCCGACCCATGG

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Artificial Sequence

SEQ ID NO: 69

CCTGAGCAGCCCCGACCCATG

DNA

Artificial Sequence

SEQ ID NO: 70

CCCTGAGCAGCCCCGACCCATGG

DNA

Artificial Sequence

SEQ ID NO: 71

CCCTGAGCAGCCCCGACCCATG

DNA

Artificial Sequence

SEQ ID NO: 72

CCCCTGAGCAGCCCCGACCCATG

DNA

Artificial Sequence

SEQ ID NO: 73

ATGTGAAGATGGGCAAGACC

SEQ ID NO: 74

ATCTCCATGTGAAGATGGGC

SEQ ID NO: 75

AACGGCCTATCTCGAGGAAT

SEQ ID NO: 76

AACATCGTCGAGCAATTCC

SEQ ID NO: 77

AATCCAACTCCTTGCCCTT

SEQ ID NO: 78

AAACCTGAGCCAGAACCTGA

SEQ ID NO: 79

AGGGCGATCTAATGAAGGGT

SEQ ID NO: 80

AGTGCACAGAAAGGACCCAC

SEQ ID NO: 81

ACACTGGTCCAGCAATGACA

SEQ ID NO: 82

TTCCTGTTGACTGAGTTGCG

SEQ ID NO: 83

CACTCTGTGGTTGGAGCAA

SEQ ID NO: 84

CAAGGCCAGGTGATGACTTT

SEQ ID NO: 85

CACACTGGTCCAGCAATGAC

SEQ ID NO: 86

CTGACACCAACCAGAGCTGA

SEQ ID NO: 87

CTCTGCCATCTGTTGGGAT

SEQ ID NO: 88

TCAAAAAGGGATCCATGCTC

SEQ ID NO: 89

TGACACCAACCAGAGCTGAG

SEQ ID NO: 90

TGATGCCTTCCTGTTGACTG

SEQ ID NO: 91

TTCCCTGTTGACTGAGTTGCG

SEQ ID NO: 92

TTCTCCAAATCGACCTTGCG

SEQ ID NO: 93

GGAGAGTTCAGGCAAAGCTG

SEQ ID NO: 94

LOCUS AY497473

DEFINITION Homo sapiens transforming growth factor, beta receptor I (activin A receptor type II-like kinase, 53kDa) (TGFBR1) gene

coding sequence in reverse complement

1 TTACATTTG ATGCCTTCCT GTTGACTGAG TTGCGATAAT GTTTCTTAA TCCGCAATGC
61 TGTAAGCCTA GCTGCTCCAT TGGCATAACCA ACATTCTCTC ATAATTAGG CCATTACTCT
121 CAAGGCTTCA CAGCTCTGCC ATCTGTTGG GATATTGGC CTTAACTTCT GTTCACAAAC
181 AACTTTCTC ATTTCTTCAA CTGATGGTC AGAAGGTACA AGATCATAAT AAGGCAGTTG
241 GTAATCTTCA TGAATTCCAC CAATGGAACA TCGTCGAGCA ATTTCCCAGA ATACTAAGCC
301 CATTGCATAG ATGTCAGCAC GTTTGAAGGA TTCAAAATGT TTCATATTAA TGGAATCATC
361 GAGAACTTCA GGGGCCATGT ACCTTTGT TCCCCTCTG TGGTTGGAG CAATATCAAT
421 GGTATCTGTG GCTGAATCAT GTCTTACTGC CAGTCCTAAG TCTGCAATAC AGCAAGTTCC
481 ATTCTTCTTT ACCAAGATAT TCTTGATTT CAAATCTCTA TGAGCAATGG CTGGCTTCC
541 TTGGGTACCA ACAATCTCCA TGTGAAGATG GGCAAGACCG CTCGCCGTGG ACAGAGCAAG
601 TTTTATCATT CCTTCCACAG TAACTGTGTA TCTGTTAAG TAATCAAAAA GGGATCCATG
661 CTCATGATAA TCTGACACCA ACCAGAGCTG AGTCCAAGTA CCATTGTCTT TATTGTCTGC
721 TGCTATAAAT CCCAGGATGT TTTCATGACG TAACATTACA GTTGATAAA TCTCTGCCTC
781 ACGGAACCAC GAACGTTCTT CTCTAGAGGA GAATATCTTA ACAGCAACTT CTTCTCCCCG
841 CCACTTTCCT CTCCAAACTT CTCCAAATCG ACCTTGCCA ATGCTTCTT GTAACACAAT
901 AGTTCTCGCA ATTGTTCTCT GAACAAGCAA TGGTAAACCT GAGCCAGAAC CTGACGTTGT
961 CATATCATAA ATTAAGTCTT TCAACGTAGT ACCCTCTGAA ATAAAAGGGC GATCTAATGA
1021 AGGGTCCTCT TCATTTGGCA CTCGATGGTG AATGACAGTG CGGTTGTGGC AGATATAGAC
1081 CATCAACATG AGTGAGATGC AGACGAAGCA CACTGGTCCA GCAATGACAG CTGCCAGTTC
1141 CACAGGACCA AGGCCAGGTG ATGACTTTAC AGTAGTTGGA AGTTCTATT TATTGCAATG
1201 GTCCTGATTG CAGCAATATG TTGTTAGTCAC AGACCCAGTT TTTGAAGAGG GTGCACATAC
1261 AAACGGCCTA TCTCGAGGAA TTAAGTCAAT TTCAGCTATA CACATGCTGT TGTGTATAAC
1321 TTTGTCTGTG GTCTCTGTGA CAGAGACAAA GCAGAGCCCA TCTGTCACAC AAGTAAAATT
1381 GTCTTTGTA CAGAGGTGGC AGAAACACTG TAACGCCGTC GCCCCCCGGGA GCAGCGCCGC

1441 CGCCGCCGCC GCCGCCGCCG CCAGCACGAG GAGGAGCAGC CGGGGACGCG GAGCAGCGAC
1501 CGCCGCCTCC AT

SEQ ID NO: 95

LOCUS AY497473

DEFINITION Homo sapiens transforming growth factor, beta receptor I (activin A receptor type II-like kinase, 53kDa) (TGFBRI) gene

mRNA in reverse complement

1 AAACACTATA GGTTTAACA TAGAGACTTT CAGCAATTCT ACTCATTCC ATTAGAAAGA
61 CACAGAAGTG GCACTTACTG GTATAGTACA ATCCCATTAA GAAGGCATGT AATGTTCTGA
121 ATAAGTGAAA GAACAATAAT AGTATACAAA ATACAATTGC ATGAATTATG TTCCTCACTA
181 CTATATGAGG TCATTTTAG ACTCAAGATA AGAGTTTAT AAGTGTAGT TTTAAGATCC
241 TGAAAAACTA TAGAACAGTA CATTCAAAGT CTGAATCAAG GAAACTCTAG TGGTCAGAA
301 TCCTCTGAGA ATGTAACAAAC CAGGAGTAAA TCACCTTCTT TAGTAATAAG ACATGTTCA
361 TTGTAATTCA GCAATCCAAC TCCTTGCCC TTAAAGATGA TCTCCAGCAC AGCAGAGTTA
421 CCTAAAGTTA AGACCAGCAA TCATCTTTT AAAAAACAAG TTTTGTAAAT AAAAAATAAA
481 GGTAGACTAC ACATTTCTG TCCTGGGAAA GAAGCGTTCA TAGTGCACAG AAAGGACCCA
541 CATGGCTGTT TCCTGGGTCC AAAGAAATCC TGGGAAGTTT TTAATTGACT TTATTACACT
601 GCTGCAAAAG GAAGCAATAT CCTCTGTTC CCTCTCAGTG AGGTAGAACAA ATTGACCTCC
661 CAAATTAAAA CCCAGGAGCA GATCTGAAGA AAAAAAGGAGA GTTCAGGCAA AGCTGTAGAA
721 TTACATTTG ATGCCTTCCT GTTGACTGAG TTGCGATAAT GTTTCTTAA TCCGCAATGC
781 TGTAAGCCTA GCTGCTCCAT TGGCATACCA ACATTCTCTC ATAATTTAG CCATTACTCT
841 CAAGGCTTCA CAGCTCTGCC ATCTGTTGG GATATTGGC CTTAACTTCT GTTCACAAAC
901 AACCTTTCTC ATTTCTTCAA CTGATGGTC AGAAGGTACA AGATCATAAT AAGGCAGTTG
961 GTAATCTTCA TGAATTCCAC CAATGGAACA TCGTCGAGCA ATTTCCCAGA ATACTAAGCC
1021 CATTGCATAG ATGTCAGCAC GTTGAAAGGA TTCAAAATGT TTCATATTAA TGGAATCATC
1081 GAGAACTTCA GGGGCCATGT ACCTTTTGT TCCCACCTCG TGGTTGGAG CAATATCAAT
1141 GGTATCTGTG GCTGAATCAT GTCTTACTGC CAGTCCTAAG TCTGCAATAC AGCAAGTTCC
1201 ATTCTTCTT ACCAAGATAT TCTTGATTT CAAATCTCTA TGAGCAATGG CTGGCTTCC
1261 TTGGGTACCA ACAATCTCCA TGTGAAGATG GGCAAGACCG CTCGCCGTGG ACAGAGCAAG
1321 TTTTATCATT CCTTCCACAG TAACTGTGTA TCTGTTAAG TAATCAAAAA GGGATCCATG
1381 CTCATGATAA TCTGACACCA ACCAGAGCTG AGTCCAAGTA CCATTGTCTT TATTGTCTGC
1441 TGCTATAAAT CCCAGGATGT TTTCATGACG TAACATTACA GTTGATAAA TCTCTGCCTC
1501 ACGGAACCAC GAACGTTCTT CTCTAGAGGA GAATATCTTA ACAGCAACTT CTTCTCCCCG
1561 CCACCTTCCT CTCCAAACTT CTCCAAATCG ACCTTGCCA ATGCTTCTT GTAACACAAT
1621 AGTTCTCGCA ATTGTTCTCT GAACAAGCAA TGGTAAACCT GAGCCAGAAC CTGACGTTGT
1681 CATATCATAA ATTAAGTCTT TCAACGTAGT ACCCTCTGAA ATAAAAGGGC GATCTAATGA
1741 AGGGTCCTCT TCATTTGGCA CTCGATGGTG AATGACAGTG CGGTTGTGGC AGATATAGAC
1801 CATCAACATG AGTGAGATGC AGACGAAGCA CACTGGTCCA GCAATGACAG CTGCCAGTTC
1861 CACAGGACCA AGGCCAGGTG ATGACTTTAC AGTAGTTGGA AGTTCTATT TATTGCAATG
1921 GTCCTGATTG CAGCAATATG TTGTAGTCAC AGACCCAGTT TTTGAAGAGG GTGCACATAC

1981 AAACGGCCTA TCTCGAGGAA TTAAGTCAAT TTCAGCTATA CACATGCTGT TGTGTATAAC
2041 TTTGTCTGTG GTCTCTGTGA CAGAGACAAA GCAGAGCCCA TCTGTCACAC AAGTAAAATT
2101 GTCTTTGTA CAGAGGTGGC AGAAACACTG TAACGCCGTC GCCCCCGGGA GCAGCGCCGC
2161 CGCCGCCGCC GCCGCCGCCG CCAGCACGAG GAGGAGCAGC CGGGGACGCG GAGCAGCGAC
2221 CGCCGCCTCC ATGGTCCCAGC CGCCACCGCC TGTGGCCCGG CCCGGCCCGG CCGCGCCGCT
2281 GCCTCACCCCC AGCAAACCTC GCCTCGCC

SEQ ID NO: 96

LOCUS AY497473

DEFINITION Homo sapiens transforming growth factor, beta receptor I
(activin A receptor type II-like kinase, 53kDa) (TGFBR1) gene

gene in reverse complement

1 CCTAGGATGG GCAGTATAAA CAACTGTGTT TTTAAATGAC AAAATTATTA CTCTAATCAA
61 ATTTATCTGC AGTTATCGAC AGTAATCATG ATCCATCCCA CCTTTAAGT TTCTTCATT
121 TTCAATATCA GCCAATTGTG GTTTATCAA CCATCCCTAG CCAAAAAAGG AATTAACTGC
181 AGTAATACGT AAAGTCAATT TAGAACTAAA CTCCAACCTT AAAAAAAATTC CCTGATTTCAT
241 CAAAAGTACA TTTTATCCAC TTAGGTAAGA AAAGTATAAT TGTAACTCAA TCTTGATTC
301 AATAAAAAAT ATCTTCAGCA AAAGTGGATA ACCTTACATG AAAGTGAGAA ATCATGTATT
361 ACAACTTAA ATCAGACAAAC TTTGGTATA TTCTAAAAT TATATAAAAA TAGTTTAGT
421 GATAAAATGC ATTGGTCCCC TGGATTCAT AAAGCAGATC TGGTAGGCTT AGAAATGGCC
481 CAAAACAAAC AACAAAAAAA AAAAAACAAA AAAAAAACCC CCACAAAAAA AGTACCTGAT
541 CACAGAATTA GAGATTATTA TTAAACAAGT AATACAACAC AACTATAAAA ACAAACCTCT
601 CAAACTCATT CATTGCTTC AATGAAATTA TGGTAACTAC AGAAAAAAAC TCAACTTTGT
661 AGTCAAAATA GTAACCTCTA AGCATTTC CCATGATGTT TTTAACACA TGAAATTGAC
721 ACTAAATTCC TATGTAATAT GTCACTCATT CTAAAATTAC TTTGGTTTA AGATCAAATA
781 GTCTTCAAA TTTATTCTAC GTATAAGAGA TAAGACATTA TCAGAATGCA ACTCATTCTAG
841 TGTAATGGC CACTTCTCT TCCTGACCAA AATACTGTAT GGGTTACTCT GACCCTGTAT
901 CGGGGAAC TG ATAGCACTAT AAAGTCGAAG TCTTGCTCA GAGCTATGAT AGTTTTCT
961 CCCCTCAGAA TAAGATCACA GTGATAAAAG GACTCGAAA ACTGTAAAAG CCCTGCAGAG
1021 ACTTCATAGC ATTGGAAAGTA AAAACCAGTT ATTATAAGCT AGTTCAACT TAATGTAAGA
1081 AGACCATGAC AAGTTGCTT TCAATATTG ACCAAAGACA TCTGTAAATG GAGAAAAACC
1141 TATACTCAGA ATGTTCTTTA GCTACCACCT CTCCCAAGTA ACAAGGGTGC ACTGAATGCA
1201 TTACATAAAAT TACATGTAAG CAGTTGCAAT GTACGCACAC ATAAGGAAAC ACTGAATTAA
1261 AAGCTGCCTT CCTCAAAGTG TAGTGAGACC TCAAAACAC ACTAGCTCTA ACCTCAGTTA
1321 CAAAGACAGA GGATCCACCG AACGTTACT CCTACATACA TAACAAATGT GACTTTCTG
1381 CACATCTGTT ATCTATTCAA ATTATTAAC ACCTTCGCCT TCCTAGAAAA AGGCCTATTA
1441 TTTAATATTA ACCATATTCT AATTTAAAT GTAATTGATG TCTTAAAAAA TAAGTCTTG
1501 AACAGATAAA GAAAAAAATG GCTTACAGAA TACCACATAA CTTTCACAA GCAGCTAGAC
1561 AGACTTCCC TTTAGAAAA ATTAGTCTGT ATGCTACAAA TATTAAAGTA CTATAAGCAT
1621 CTAGAGTGCC ACTTGATTT TTAAAGTAT ACTTAGACAA CACAATTAAA ATGCAAAAGC
1681 TTGATGTGAG AATATTCAA CATGACCATG CTAATAATT ACTAATTAAAT GCCACTTCAT

1741 TTCTTAGTG GCTTAAGCAC ATTTGGTAG GAGAAAGGCA TTTTCAGAAT AGAATACCAA
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46441 GCCCGGCAGC CGCCGAGCCG CGCGCGCCA CGAGAGCCCG GCCCAGGGCCC CGGCGCCGCC
46501 ACCTGCGCCC CGGGCCCCGC GCCATGTTTG AGAAAGAGCA GGAGCGAGCC AGAGGCCGGG
46561 TCCGGCCCGC GCGCCCCGCA GTCGCCCGCC CGCCGCGCCG CCGCTCACCC GTCGCCCCCG

46621 GGAGCAGCGC CGCCGCCGCC GCCGCCGCC CGGCCAGCAC GAGGAGGAGC AGCCGGGGAC
46681 GCGGAGCAGC GACCGCCGCC TCCATGGTCC CGCCGCCACC GCCTGTGGCC CGGCCCGGCC
46741 CGGCCGCCGC GCTGCCTCAC CCCAGCAAAC CTCGCCTCGC CCCACCTCCC TAGCCGCCGC
46801 GGCGGCCTCG CTCCGGCCCT TTGTAACTGC TCGGAGGACG CGCGTCCATT GGCTGCCGG
46861 CTCCCGCCGG CCCCGCCTCC CCGCCGCCGC GAGCTGCCAA GCGGGACCCA GCCGGGAGCC
46921 CCGCCTGCAGG GCCCGCCAGG CAGCCAATCC GCAGCCGCAG GCGCCGGTTT CTGGCCACGC
46981 CCCACGCTCC CGGGGGGCTG GGCGGCCAGA CCCCAGCCCC GGCGCGATCG GCTCCCGGCT
47041 CCGAGAGGCC GCGTGGGGGC GGGGTCTGCC AGCCCCAGCA CCGCTCAGCC GCTAGCCCCG
47101 GAGGGCCGGG TAGAGCGATG GGTGTGTCTG TGTGAGTCTC TTTCGGAAAA AGGCTGTGGC
47161 CGTCGACGC TCTTTCTTC TAACCTCCTC TAGGCAGCGA AGATCTGGTA CTGCCTCAGC
47221 CCCACCCCTGA CCCCATTACAG AGCTCCGCAT TGGCAATCCC AGCACATGCC ACCCAGATCC
47281 GCTGCAGCGC CAGGCCCTG CATCCACTTC AACTCCCCCT CTCCAAGTCC ACGCATCAAC
47341 TTCAGCATTG CCCCAGAGAT CCCTCCTCAG TGCCCTTCAC ATGCGACTCA CTCTCCTATT
47401 TCCTCCTGCT CCCAGCGGTC ACCCCAAACC CAGCTCCCCC ACCCAGTTCC AAACCCAGAA
47461 AGTCCTCAGA TCCCAGCGTC AGACCCAGAT CCTGAGCCCA AACACACCCC CAAATAGCCT
47521 CCCGCCTCCC TCCAGCACAG ATCCAGGATG GGGATCCAAG CCGCACTCCT CTCAAACCCC
47581 TTCCCGATCC AGGTTGGAAA GGGAGGATCC CCACCCCAAC CCCTCAAAGG AGGGGTTCCC
47641 CCTCTTTAGC ACCCAGCTCC CGCGGGGGCGG AGGGGGAGCA GTCATCATT A CTTGAGCTG
47701 TGTCTGACAC TGCTGTATAA TAATCCTGAG GTGTCACTAA AAACCCAAAT AAACCTACG
47761 TTTTCATTCT GAATTCTAA TTTACTGCGA GACGCTCCAC CCACCTTCCC TCTGCGACGC
47821 CAAAATGAGC TCCAGATTTG TAATTCTCC TGTCAGACTA GTCCCTTCTT CAGACACAAA
47881 CACAGCCAAG GAGGCTGTTA CGTAGAACAG AGAATATTT TCCGCAGAAC CTTTAAGGA
47941 GGAAATTTA TTTCTGTGT CATTGAGCT TAGAATTAAA TAAACCTTGA TAGCAGGAAT
48001 CAGAATGGTT CTGATTAATG CCAATTTGTG CACTACTTA GAGTCACTGG GTGAGCATT
48061 TCCCCACCTG TGAAGGCTTC CTTCCGGGAG ATGAAAAGGG AAAAGGCGTG GATATTGGAG
48121 CTGGGATCTG AGACCCCGCGC GCTTTCTCCC TCCTGTCTAG GACACTTACT TAAGCAAGTC
48181 AGCTAACCTT TCTGAACCTC CAACAGGCTC AGAGGCCTGC AACTGCTCT ACACCGCAGC
48241 TCCAGCAGCC TCAGCAGCAA ACTCCCCTCT TGCTCAGGCT GAGGGATTCC AGAGAGATGG
48301 CTTCTGGCAG GCTCCAGTCC CAATTGCC TCTCAGTTCC AGTTCTTCA ATGGAGATTT
48361 GGTGGACTTT GTGCCACCTG AGGTCCCTAG ACTGCCTTT TGCTGCTCTA TTTGCAATGT
48421 CTTTCATAAG ATAAGAGCTA ATGAGATTT TATTGTATGA ATGAATGAAT GATCTCAGAA
48481 AGTGAATGTT TTTCTCTGGC CATTAGCTTC TTCATCTCTA AAATGTAAAT AATAATAGTA
48541 CCTACCCCGT TTATAAATAA AAGGGATTGA ATCACTTGTG TTGGGCCCT TAAATGACGT
48601 AATGTGTCTG GTACAGTGTG TGAAACATAG TAAATATTTA GCTAATGCCA TTTCTTGCC
48661 CATCCCCCTTC CAGCTCTGTA TGATTCTAAT CAGCATGTAT GTTCATGTCA GTCTGTGCCT
48721 TTGACGTACG GAGCCTAGAT TAATCAGTGT TAATCACACC TCCAGTTCT TAACA